

Figure.1

### 1st Amino Acid Sequence

File Name : clone 106.full.seq.protein  
Sequence Size : 222

### 2nd Amino Acid Sequence

File Name : TSG.protein  
Sequence Size : 249  
[33.3% / 216 aa]

1' MKSHYIVLALASLTFLCLPVSQSCNKALCASDVSKCLIQELCQCRPGEGNCPCCKECM  
\*.\*. .... .\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. ....  
1" MQLLCYFVILFVGIAFWSSLANDDGCNEVCGSVSKCLITQSCQCKLND—CHCKDCL  
60' LCLGALWDECCDCVGMCNPRNYSDTPPPTSKSTVEELHEPIPSLFRALTEGDTQLNWNIVS  
\*\*\*.\*. \*\*\*.\*. \*\*\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. ....  
59" NCLGELYIECCGCLDMCPKHKDVPLSLTPRSEIGDI-EGVPELFDTLTAEDDE-GWSTIR  
120' FPVAEELSHHENLVSFLETVNQLHHQNVSPSNNVHAPFPSDKERMCTVVFDDCMSIHQ  
\*.. .... .\*. .... \*.. .... \*.. .... \*.. .... \*.. .... \*.. .... \*.. ....  
117" FSMRAGFKQR—V——QGGASGDAGNGNGNGNAGSAGVTLCTVIYVNSCI RANK  
180' CKISCESMGASKYRWFHNACCECIGPECIDYGSKTVKCMNCMF  
\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. .... \*.\*. ....  
166" CRQQCESMGASSYRWFHDGCCCEVGENCLNYGINESRCRGCPEDQDQLLTADTVPAEAEQ